PC-0039 US

<110> Chen, Huei-Mei
Honchell, Cynthia D.
Tang, Y. Tom

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acagaaagga accgagtgat tgtggatgct accactggat tgatcccttt gaccagtgta 600
cccacatcag caaaagaaat gaccacaaag cttggggtta cag 643

<210> 6
<211> 554
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 7437161H1

<400> 6
tgtacccaca tcagcaaaag aaatgaccac aaagcttggc gttacagcag agtacagccc 60
agcttcacgt tccctcggaa catctccttc tccccaaacc acagttgttt ccacggctga 120
agacttggct cccaaatctg ccacctttgc tgttcagagc agcacacagt caccaacaac 180
actgtcctct tcagcctcag tcaacagctg tgctgtgaac ccttgtcttc acaatggcga 240
atgcgtcgca gacaacacca gccgtggcta ccactgcagg tgcccgcctt cctggcaagg 300
ggatgattgc agtgtggatg tgaatgagtg cctgtcgaac ccctgcccat ccacagccac 360
gtgcaacaat actcagggat cctttatctg caaatgcccg gttgggtacc agttggaaaa 420
agggatatgc aatttggtta gaaccttcgt gacagagttt aaattaaaga gaacttttct 480
taatacaact gtggaaaaac attcagacct acaagaagtt gaaaatgaga tcaccaaaac 540
gttaaatatg tgtt 554

<210> 7
<211> 571
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 71247228V1

<400> 7
gatcaccaaa acgttaaata tgtgtttttc agcgttacct agttacatcc gatctacagt 60
tcacgcctct agggagtcca acgcggtggt gatctcactg caaacaacct tttccctggc 120

ctccaatgtg acgctatttg acctggctga taggatgcag aaatgtgtca actcctgcaa 180
ggtcctctgc tgaggtctgc cagctcttgg gatctcagag gcggatcttt agagcgggca 240
gcttgtgcaa gcggaagagt cccgaatgtg acaaagacac ctccatctgc actgacctgg 300
acggcgttgc cctgtgccag tgcaagtcgg gatactttca gttcaacaag atggaccact 360
cctgccgagc atgtgaagat ggatataggc ttgaaaatga aacctgcatg agttgcccat 420
ttggccttgg tggtctcaac tgtggaaacc cctatcagct tatcactgtg gtgatcgcag 480
ccgcgggagg tgggctcctg ctcatcctag gcatcgcact gattgttacc tgttgcagaa 540
agaataaaaa tgacataagc aaactcatct t 571

<210> 8
<211> 433
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 6475676H1

<400> 8
tgaaacttgc atgagttgtc cattcagcct tggtggtctc aactgtggaa acccctatca 60
gcttatcact gtggtgatcg cagccgcggg aggtgggctc ctgctcatcc taggcatcgc 120
actgattgtt acctgttgca gaaagaataa aaatgacata agcaaactca tcttcaaaag 180
tggagatttc caaatgtccc cgtatgctga ataccccaaa aatcctcgct cacaagaatg 240
gggccgagaa gctattgaaa tgcatgagaa tggaagtacc aaaaacctcc tccagatgac 300
ggatgtgtac tactcgccta caagtgtaag gaatccagaa cttgaacgaa acggactcta 360
cccgggctac actggactgc caggatcacg ggattcttgc attttccccg gacagtataa 420
accgtctttc atc 433

<210> 9
<211> 538
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 7735769H1

<400> 9
ggggccgaga agctattgaa atgcatgaga atggaagtac caaaaacctc ctccagatga 60
cggatgtgta ctactcgcct acaagtgtaa ggaatccaga acttgaacga aacggactct 120
acccggccta cactggactg ccaggatcac ggcattcttg catttcccc ggacagtata 180
acccgtcttt catcagtgat gaaagcagaa gaagagacta cttttaagtc caggagagag 240
agggactcat tgctctgagc cagtcacctg ggacctctgc tcagaggacc gcaccaggag 300
gctgcgccca ggatttgtcg ggagccacgc tgagtggcaa gcaggaacga gggacaggca 360
tgcggggcgt gaccacagtg gaggagacag gtggatgtgg aaccacaggc tgctcattca 420
gcacctttgt tgttactgtg aacgtgaatg tgggccagta tcaagagagt ctctctgagt 480
gactgcacca tggcactggc accagggcga ctattagcca gggcagacca ctagactt 538

<210> 10
<211> 567
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 7180688H1

<400> 10
ctagacttca gtgcaggacc tggttttccc ttcgtttgca ctttagtaaa ttgggtggga 60
ggtttccttt tggatctgtt ttgagactgt tccagaaaga aggcttcctt tcccgagaca 120
cttccatagg cagcaatttg gtgattcatt tgcagcaaaa tactggcttg ttaattattt 180

tcctgcccag cgcctgcgtg ctaaacaaca gatgaggatg agcgtaccac tgaagtctga 240
agatgtcgcc attgaacgga cagtgttttc atatgtttct aggttgtctt atgctacagt 300
ttccaagcca gcccccacag tgaggaaatg tgtgaggcac cgcacacaac tgcaatgtgt 360
tttttaagtc aaggtgacac atgtatttaa gattttttt taaaatctct ttgcagttaa 420
atctcacttt ttcaaacaag cctggatcag ggcaaaacaa cttatatttg gttttagctg 480
gaggctcagc aggcagattg caggcagggg ggcacttttc atccatgaga ggccagcctg 540
gggcctggga ctctgatcac cattgtg 567

<210> 11
<211> 600
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 70650868V1

<400> 11
ctcacttcat ccaaaaccag gatgccccaa agaaggccaa taaacacagt tccccaggtg 60
gcaattaaag acaccgagta ctggatgtct ccctggcagg acccacatca caggcataat 120
aaataagatg agtggaactt ccttcccgaa ggtcaaccct cagttcctcg accaaccgga 180
agtcttcagt tctcccacac tgactggaag tataaccacg tttctggagg gtgcgacaca 240
gccatatgaa gaggtacaaa tgactgggtg agaaaaaaaa gttatttctt cagccgaata 300
aacagctttg agtggttgaa agtttacatg ggtttgtgg acatgagatt ctgggtacaa 360
agtgctgcag tagccggtga gcaaactcat gtgtggctcc atctcggctc cctgttcttc 420
ctcaggaatc cacacagctt cccaaagcac tgttgatgca ggaaatctaa cctggctatt 480
cagcccatcc ctctaaccac atccagctgc aggggctcaa caagctgctt tcctagagtg 540
gtgaaacctg cgttcagttt gacattttct cctccataag caggttgctc tggcctccac 600

<210> 12
<211> 371
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 2359874T6

<400> 12
gaagaaacaa ccatgcccgg ggcagaactg tccccagtgg ctgcaccatt tccagaaatg 60
tgaagcggga ctccccgctt cagggtggag acaattcttt tacctctgta ttcccctcac 120
ttcatccaaa accaggatgc cccaaagaag gccaataaac acagttcccc aggtggcaat 180
taaagacacc gagtactgga tgtctccctg gcaggaccca catcacaggc ataataaata 240
agatgagtgg aacttccttc ccgaagtcaa ccctcagttc ctcgaccaac cggaagtctt 300
cagttctccc acactgactg gaagtataac cacgtttctg gagggtgcga cacagccata 360
tgaaggaatt c 371

<210> 13
<211> 399
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 2359874R6

<400> 13
cttcatatgg ctgtgtcgca ccctccagaa acgtggttat acttccagtc agtgtgggag 60
aactgaagac ttccggttgg tcgaggaact gagggttgac cttcgggaag gaagttccac 120
tcatcttatt tattatgcct gtgatgtggg tcctgccagg gagacatcca gtactcggtg 180
tctttaattg ccacctgggg aactgtgttt attggccttc tttggggcat cctggttttg 240

gatgaagtga ggggaataca gaggtaaaag aattgtctcc accctgaagc ggggagtccc 300
gcttcacatt tctggaaatg gtgcagccac tggggacagt tctgccccgg gcatggttgt 360
ttcttcaagg tcctctaaat ataatcccta ttcttacat 399

<210> 14
<211> 595
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 70650365V1

<400> 14
tttggggcat cctggttttg gatgaagtga ggggaataca gaggtaaaag aattgtctcc 60
accctgaagc ggggagtccc gcttcacatt tctggaaatg gtgcagccac tggggacagt 120
tctgccccgg gcatggttgt ttcttcaagg tcctctaaat ataatcccta ttcttacata 180
atcctgtggc ctgatggttt taagcaagaa ctcctgtgtc ccatggtctc caccactcac 240
catcaccctg ctgtagcaag agtcctagtc aggggaggtg cattttagta gttaaatggc 300
acttatccat gagataaata aaaggagaac tgtttttatc agtggaggct aacctaaaat 360
ttcaaagtgt cgccttttgg aaatctgggg cctctctctc tgtagaacca atggcccttg 420
gtggctcacg gcctcgcacc ctaactggag agttctgagc tcctgcagct cacctgagcc 480
cacagactag gcttcttggc tccttccgca gcaggctggt tcaccccaga acccgcagct 540
gtgggaagag ccatgtaggg aggctaatcc caggcataca cttccactgc cttca 595

<210> 15
<211> 549
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 1241344R6

<220>
<221> unsure
<222> 442, 460, 515
<223> a, t, c, g, or other

<400> 15
acctaactgg agagttctga gctcctgcag ctcacctgag cccacagact aggcttcttg 60
gctccttccg cagcatgcct gctcaccccc agaacccgca gctgtgggaa gagccatgta 120
gggaggctat tcccaggcat acacttccac tgccttcagc tgacgtcaca gctgacaaat 180
catctcctct atcggagcca gaagacttca gctccacaaa atgaagtgtt ctgtcctgaa 240
aacattcttg ggaagaatcc caacatcgag aaaacggtgt cctgtgagtt ccaacaatgc 300
ttcttgttca tgggtttctt ccgtatggag tggattaaga gtgttttatt ttgttgttct 360
aactgagaaa aaaaggaggc acccacaagg ttgaggtcac acagtctcca cagtttccag 420
gaggcgtttg ggggtgggga angcacctcc agagcatgan ggctctaagg ggacatgagt 480
aaagcatgtc tgtgacccag tgaggaaagg gagangccag ctgcactcct gcaacggggg 540
ttcctagct 549

<210> 16
<211> 272
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 008938H1

<220>

<221> unsure
<222> 75, 106, 112, 163, 167, 192, 252
<223> a, t, c, g, or other

<400> 16
ggagaggcca gctgcactcc tgcacggggt tcctagctgc agaagggtcc cgcctaggcc 60
gaggggaaac acctnatagc agaagaggcc tggatgcaca cctggnacgc cnaggctctc 120
cgcccagaca cagtgctcca tgtcaacccc tgcacctggg gtntgtnatt cacgtgcaca 180
gatgccacaa tnctgcacca atatcccaca gatgggggaa ggtgagagga aggggcaagt 240
aatgtgtacc tnctcaagag atgcttaaac ct 272

<210> 17
<211> 424
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 2580841F6

<220>
<221> unsure
<222> 162, 251
<223> a, t, c, g, or other

<400> 17
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aaggccctgg agtctgtttc tttaggcgga tgaactgaca tgctcctacc atgaccaggc 120
tctgggcaag gctcctcaca gtatccttga gaggtgggca tngaagtgcc catttctcag 180
gtacagaaac cttcagagag gataaatagc ttgccctgta gaagcaggac tgaaaccctt 240
gtccgcctga ntcccccagc tactctgccc actgtagccc cctgccttac tgtcctggca 300
caccctctac catcctgtat accttaaata tcaaagaggg caagagagaa agggctttaa 360
agataagtta ttttttaag gaaccttaat attatttta agaagtaacc aaattagtga 420
cgtg 424

<210> 18
<211> 430
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 70621193V1

<400> 18
cctggtacac ccctcaccat cctgtatacc ttaaatatca aagagggcaa gagagaaagg 60
gctttaaaga taagttattt ttttaaggaa ccttaatatt atttttaaga agtaaccaaa 120
ttagtgacgt gaaatgcaaa aaaaaaaaaa aaaaatgtct gactaccctt ttggaaaagt 180
gtgcttccag attggctttt ttatagtgta attctttaga cacttggtca ttaagaaaaa 240
tagtggcggg ctggtgcttc agcaagaagc acacgggcac ggtggcttgg gatataggag 300
gtggaaggca aggaccgggt gtttctggac aggtggcggc cagacttaca cttccatctg 360
gagagctggt ggctttggtc ccctgggtag ggccatgggt tccccactat tactgggaag 420
ctatagggtg 430

<210> 19
<211> 957
<212> PRT
<213> Homo sapiens

<220>
<221> misc_feature

PC-0039 US

<223> Genbank ID No: g2853301

<400> 19

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Ile Thr Ile Thr Glu Thr Thr Ser His Ser Thr Pro Ser Tyr Thr
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Thr Ser Ile Thr Thr Thr Glu Thr Pro Ser His Ser Thr Pro Ser
             20                  25                  30
Tyr Thr Thr Ser Ile Thr Thr Thr Glu Thr Pro Ser His Ser Thr
             35                  40                  45
Pro Ser Phe Thr Ser Ser Ile Thr Thr Thr Glu Thr Thr Ser His
             50                  55                  60
Ser Thr Pro Ser Phe Thr Ser Ser Ile Arg Thr Thr Glu Thr Thr
             65                  70                  75
Ser Tyr Ser Thr Pro Ser Phe Thr Ser Ser Asn Thr Ile Thr Glu
             80                  85                  90
Thr Thr Ser His Ser Thr Pro Ser Tyr Ile Thr Ser Ile Thr Thr
             95                 100                 105
Thr Glu Thr Pro Ser Ser Ser Thr Pro Ser Phe Ser Ser Ser Ile
            110                 115                 120
Thr Thr Thr Glu Thr Thr Ser His Ser Thr Pro Gly Phe Thr Ser
            125                 130                 135
Ser Ile Thr Thr Thr Glu Thr Thr Ser His Ser Thr Pro Ser Phe
            140                 145                 150
Thr Ser Ser Ile Thr Thr Thr Glu Thr Thr Ser His Asp Thr Pro
            155                 160                 165
Ser Phe Thr Ser Ser Ile Thr Thr Ser Glu Thr Pro Ser His Ser
            170                 175                 180
Thr Pro Ser Ser Thr Ser Leu Ile Thr Thr Thr Lys Thr Thr Ser
            185                 190                 195
His Ser Thr Pro Ser Phe Thr Ser Ser Ile Thr Thr Thr Glu Thr
            200                 205                 210
Thr Ser His Ser Ala Arg Ser Phe Thr Ser Ser Ile Thr Thr Thr
            215                 220                 225
Glu Thr Thr Ser His Asn Thr Arg Ser Phe Thr Ser Ser Ile Thr
            230                 235                 240
Thr Thr Glu Thr Asn Ser His Ser Thr Thr Ser Phe Thr Ser Ser
            245                 250                 255
Ile Thr Thr Thr Glu Thr Thr Ser His Ser Thr Pro Ser Phe Ser
            260                 265                 270
Ser Ser Ile Thr Thr Thr Glu Thr Pro Leu His Ser Thr Pro Gly
            275                 280                 285
Leu Pro Ser Trp Val Thr Thr Thr Lys Thr Thr Ser His Ile Thr
            290                 295                 300
Pro Gly Leu Thr Ser Ser Ile Thr Thr Thr Glu Thr Thr Ser His
            305                 310                 315
Ser Thr Pro Gly Phe Thr Ser Ser Ile Thr Thr Thr Glu Thr Thr
            320                 325                 330
Ser Glu Ser Thr Pro Ser Leu Ser Ser Ser Thr Ile Tyr Ser Thr
            335                 340                 345
Val Ser Thr Ser Thr Thr Ala Ile Thr Ser His Phe Thr Thr Ser
            350                 355                 360
Glu Thr Ala Val Thr Pro Thr Pro Val Thr Pro Ser Ser Leu Ser
            365                 370                 375
Thr Asp Ile Pro Thr Thr Ser Leu Arg Thr Leu Thr Pro Ser Ser
            380                 385                 390
Val Gly Thr Ser Thr Ser Leu Thr Thr Thr Thr Asp Phe Pro Ser
            395                 400                 405
Ile Pro Thr Asp Ile Ser Thr Leu Pro Thr Arg Thr His Ile Ile
            410                 415                 420
Ser Ser Ser Pro Ser Ile Gln Ser Thr Glu Thr Ser Ser Leu Val
            425                 430                 435
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Gly Thr Thr Ser Pro Thr Met Ser Thr Val Arg Met Thr Leu Arg
440 445 450
Ile Thr Glu Asn Thr Pro Ile Ser Ser Phe Ser Thr Ser Ile Val
455 460 465
Val Ile Pro Glu Thr Pro Thr Gln Thr Pro Pro Val Leu Thr Ser
470 475 480
Ala Thr Gly Thr Gln Thr Ser Pro Ala Pro Thr Thr Val Thr Phe
485 490 495
Gly Ser Thr Asp Ser Ser Thr Ser Thr Leu His Thr Leu Thr Pro
500 505 510
Ser Thr Ala Leu Ser Thr Ile Val Ser Thr Ser Gln Val Pro Ile
515 520 525
Pro Ser Thr His Ser Ser Thr Leu Gln Thr Thr Pro Ser Thr Pro
530 535 540
Ser Leu Gln Thr Ser Leu Thr Ser Thr Ser Glu Phe Thr Thr Glu
545 550 555
Ser Phe Thr Arg Gly Ser Thr Ser Thr Asn Ala Ile Leu Thr Ser
560 565 570
Phe Ser Thr Ile Ile Trp Ser Ser Thr Pro Thr Ile Ile Met Ser
575 580 585
Ser Ser Pro Ser Ser Ala Ser Ile Thr Pro Val Phe Ser Thr Thr
590 595 600
Ile His Ser Val Pro Ser Ser Pro Tyr Ile Phe Ser Thr Glu Asn
605 610 615
Val Gly Ser Ala Ser Ile Thr Gly Phe Pro Ser Leu Ser Ser Ser
620 625 630
Ala Thr Thr Ser Thr Ser Ser Thr Ser Ser Ser Leu Thr Thr Ala
635 640 645
Leu Thr Glu Ile Thr Pro Phe Ser Tyr Ile Ser Leu Pro Ser Thr
650 655 660
Thr Pro Cys Pro Gly Thr Ile Thr Ile Thr Ile Val Pro Ala Ser
665 670 675
Pro Thr Asp Pro Cys Val Glu Met Asp Pro Ser Thr Glu Ala Thr
680 685 690
Ser Pro Pro Thr Thr Pro Leu Thr Val Phe Pro Phe Thr Thr Glu
695 700 705
Met Val Thr Cys Pro Thr Ser Ile Ser Ile Gln Thr Thr Leu Thr
710 715 720
Thr Tyr Met Asp Thr Ser Ser Met Met Pro Glu Ser Glu Ser Ser
725 730 735
Ile Ser Pro Asn Ala Ser Ser Ser Thr Gly Thr Gly Thr Val Pro
740 745 750
Thr Asn Thr Val Phe Thr Ser Thr Arg Leu Pro Thr Ser Glu Thr
755 760 765
Trp Leu Ser Asn Ser Ser Val Ile Pro Leu Pro Leu Pro Gly Val
770 775 780
Ser Thr Ile Pro Leu Thr Met Lys Pro Ser Ser Ser Leu Pro Thr
785 790 795
Ile Leu Arg Thr Ser Ser Lys Ser Thr His Pro Ser Pro Pro Thr
800 805 810
Thr Arg Thr Ser Glu Thr Pro Val Ala Thr Thr Gln Thr Pro Thr
815 820 825
Thr Leu Thr Ser Arg Arg Thr Thr Arg Ile Thr Ser Gln Met Thr
830 835 840
Thr Gln Ser Thr Leu Thr Thr Thr Ala Gly Thr Cys Asp Asn Gly
845 850 855
Gly Thr Trp Glu Gln Gly Gln Cys Ala Cys Leu Pro Gly Phe Ser
860 865 870
Gly Asp Arg Cys Gln Leu Gln Thr Arg Cys Gln Asn Gly Gly Gln
875 880 885
Trp Asp Gly Leu Lys Cys Gln Cys Pro Ser Thr Phe Tyr Gly Ser

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                  890                 895                 900
Ser Cys Glu Phe Ala Val Glu Gln Val Asp Leu Asp Ala Glu Asp
                  905                 910                 915
Phe Cys Arg His Ala Gly Leu His Leu Gln Gly Cys Gly Asp Pro
                  920                 925                 930
Val Pro Glu Glu Trp Gln His Arg Gly Gly Leu Pro Gly Pro Ala
                  935                 940                 945
Gly Asp Ala Leu Gln Pro Pro Ala Gly Glu Arg Val
                  950                 955

<210> 20
<211> 528
<212> PRT
<213> Sus scrofa

<220>
<221> misc_feature
<223> Genbank ID No: g915208

<400> 20
Pro Ile Ser Val Gln Pro Ser Ser Ser Ser Ser Ser Pro Thr Thr
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Ser Thr Thr Ser Val Gln Ser Ser Ser Ser Ser Ser Val Pro Ile
                 20                  25                  30
Pro Ser Thr Thr Ser Val Gln Pro Ser Ser Ser Gly Ser Ala Pro
                 35                  40                  45
Thr Thr Ser Ala Thr Ser Val Gln Thr Ser Ser Ser Ser Ser Pro
                 50                  55                  60
Pro Ile Ser Ser Thr Ile Ser Val Gln Thr Ser Ser Ser Ser Ser
                 65                  70                  75
Val Pro Thr Thr Ser Thr Thr Ser Val Gln Pro Ser Ser Ser Ser
                 80                  85                  90
Ser Ala Pro Thr Thr Arg Ala Thr Ser Val Gln Ser Ser Ser Ser
                 95                 100                 105
Ser Ser Ala Pro Ile Ser Ser Thr Thr Ser Val Gln Pro Ser Ser
                110                 115                 120
Ser Gly Ser Val Pro Thr Thr Ser Ala Thr Ser Val Gln Ser Ser
                125                 130                 135
Ser Ser Ser Ser Ala Pro Thr Thr Ser Ala Thr Ser Val Gln Pro
                140                 145                 150
Ser Ser Ser Ser Ser Pro Pro Ile Ser Ser Thr Val Ser Val Gln
                155                 160                 165
Pro Ser Ser Ser Ser Ser Ala Pro Thr Thr Ser Ala Thr Ser Val
                170                 175                 180
Gln Pro Ser Ser Ser Ser Ser Pro Pro Ile Ser Ser Thr Val Ser
                185                 190                 195
Val Gln Thr Ser Ser Ser Ser Ser Val Pro Thr Thr Ser Thr Thr
                200                 205                 210
Ser Val Gln Pro Ser Ser Ser Ser Ser Val Pro Thr Thr Ser Ala
                215                 220                 225
Thr Ser Val Arg Ser Ser Ser Ser Ser Ser Thr Pro Ile Pro Ser
                230                 235                 240
Thr Thr Ser Val Gln Pro Ser Ser Ser Ser Ser Ala Pro Thr Thr
                245                 250                 255
Ser Ala Thr Ser Val Gln Pro Ser Ser Ser Ser Ser Thr Pro Ile
                260                 265                 270
Pro Ser Thr Thr Ser Val Gln Pro Ser Ser Ser Ser Ser Ala Pro
                275                 280                 285
Thr Thr Ser Ala Thr Ser Val Gln Pro Ser Ser Ser Ser Ser Pro
                290                 295                 300
Pro Ile Ser Ser Thr Ile Ser Val Gln Pro Ser Ser Ser Ser Ser
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                305                 310                 315
Ser Pro Thr Thr Ser Thr Thr Ser Val Gln Pro Ser Ser Ser Gly
                320                 325                 330
Ser Ala Pro Thr Thr Ser Ala Thr Ser Val Gln Pro Ser Ser Ser
                335                 340                 345
Ser Ser Pro Pro Ile Ser Ser Thr Ile Ser Val Gln Pro Ser Ser
                350                 355                 360
Ser Ser Ser Ser Pro Thr Thr Ser Thr Thr Ser Val Gln Pro Ser
                365                 370                 375
Ser Ser Gly Ser Ala Pro Thr Thr Ser Ala Thr Ser Val Gln Pro
                380                 385                 390
Ser Ser Ser Ser Ser Val Pro Thr Thr Ser Ala Thr Ser Val Arg
                395                 400                 405
Ser Ser Ser Ser Ser Ser Thr Pro Ile Pro Thr Thr Thr Ser Val
                410                 415                 420
Gln Pro Ser Ser Ser Ser Ser Val Pro Thr Thr Ser Ala Thr Ser
                425                 430                 435
Val Gln Thr Ser Ser Ser Ser Ser Thr Pro Ile Pro Ser Thr Thr
                440                 445                 450
Ser Val Gln Pro Ser Ser Ser Ser Ser Ala Pro Thr Thr Ser Ala
                455                 460                 465
Thr Ser Val Gln Pro Ser Ser Ser Ser Ser Pro Pro Ile Ser Ser
                470                 475                 480
Thr Ile Ser Val Gln Pro Ser Ser Ser Ser Ser Ser Pro Thr Thr
                485                 490                 495
Ser Thr Thr Ser Val Gln Pro Ser Ser Ser Gly Ser Ala Pro Thr
                500                 505                 510
Thr Ser Ala Thr Ser Val Gln Pro Ser Ser Ser Ser Ser Pro Pro
                515                 520                 525
Ile Ser Ser
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